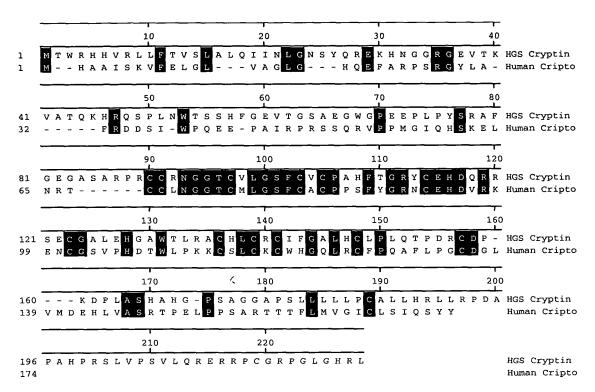
## FIG. 1

				CTG Leu 10						48
				TAT Tyr				AAC , Asn		96
				GCC Ala					1	44
				TTC Phe					1	.92
				CTC Leu					. 2	40
				TGC Cys 90					2	88
 	 	 	 	 CCG Pro	 -	_	 		3	36
				GAA Glu					3	84
				CTC Leu					4	32
				CCT Pro					4	180
				CCG Pro 170					5	28
				CTC Leu					5	576
				CTG Leu					6	524
				GGA Gly				TAA *	6	572

AMINO ACID ALIGNMENT BETWEEN CRIPTO AND CRIPTIN

FIG. 2



Decoration 'Decoration #1': Shade (with solid black) residues that match the consensus named 'Consensus #1' exactly.